

Fig. 1**SEQ ID NO: 1**

GGGCCGGGCAGGGCCGGGGCGTGGGCCGGCAGGAAGATGGCGAACGTG
GGGCTGCAGTTCCAGGCGAGCGCGGGGGACTCGGACCCACAGAGCCGG
CCCCTGCTGCTGCTCGGGCAGCTGCACCACCTGCACCGCGTGCCCTGG
AGCCACGTCCGCGGGAAGCTGCAGCCCCGGGTACCGAGGAGCTCTGG
CAGGCTGCCCTGAGCACGCTCAACCCCAACCCACGGACAGCTGTCCC
CTCTACCTAACTACGCCACCGTGGCTGCCCTGCCCTGCAGGGTGAGCC
GGCACAACAGCCCCTCGGCCGCCCACTTCATCACGCGGCTGGTGCGGA
CCTGCCTGCCGCCCGGAGCGCATCGCTGCATTGTGATGGTCTGCGAGC
AGCCAGAGGTCTTTGCTTCCGCCTGTGCCCTGGCCCGGGCCTTCCCGC
TGTTACCCACCGCTCAGGTGCCTCTCGGCGCTTGGAGAAGAAGACGG
TCACCGTGGAGTTTTTCTGGTGGGACAAGACAACGGGCCGGTGGAGG
TGTCACATTCAGTGCTTAGCGAATGCCACAGACGGCGTGCGGCTAG
CAGCCCGCATCGTGGACACACCCTGCAATGAGATGAACACCGACACCT
TCCTCGAGGAGATTAACAAAGCTGGAAAGGAGCTGGGGATCATCCCAA
CCATCATCCGGGATGAGGAACTGAAGACGAGAGGATTTGGAGGAATCT
ATGGGGTTGGCAAAGCCGCCCTGCATCCCCCAGCCCTGGCCGTCTCA
GCCACACCCAGATGGAGCCACGCAGACCATCGCCTGGGTGGGCAAAG
GCATCGTCTATGACACTGGAGGCCTCAGCATCAAAGGGAAGACTACCA
TGCCGGGGATGAAGCGAGACTGCGGGGGTGCTGCGGCCGTCTGGGGG
CCTTCAGAGCCGCAATCAAGCAGGGTTTCAAAGACAACCTCCACGCTG
TGTTCTGCTTGGCTGAGAACTCGGTGGGGCCCAATGCGACAGGGCCAG
ATGACATCCACCTGCTGTACTCAGGGAAGACGGTGGAAATCAACAACA
CGGATGCCGAGGGCAGGCTGGTGCTGGCAGATGGCGTGTCTATGCTT
GCAAGGACCTGGGGGCCGACATCATCCTGGACATGGCCACCCTGACCG
GGGCTCAGGGCATTGCCACAGGGAAGTACCACGCCGCGGTGCTCACCA
ACAGCGCTGAGTGGGAGGCCGCTGTGTGAAGGCGGGCAGGAAGTGTG
GGGACCTGGTGCACCCGCTGGTCTACTGCCCCGAGCTGCACTTCAGCG
AGTTCACCTCAGCTGTGGCGGACATGAAGAACTCAGTGGCGGACCGAG

ACAACAGCCCCAGCTCCTGTGCTGGCCTCTTCATCGCCTCACACATCG
GCTTCGACTGGCCCGGAGTCTGGGTCCACCTGGACATTGCTGCACCGG
TGCATGCTGGTGAGCGAGCCACAGGCTTCGGTGTGGCCCTCCTGCTGG
CGCTCTTCGGCCGTGCCTCTGAGGACCCTCTGCTGAACCTGGTGTCCC
CACTGGGCTGTGAGGTGGATGTCGAGGAGGGGGACCTGGGGAGGGACT
CCAAGAGACGCAGGCTTGTGTGAGCCTCCTGCCTCGGCCCTGACAAAC
GGGGATCTTTTACCTCACTTTGCACTGATTAATTTTAAGCAATTGAAA
GATTGCCCTTC

Fig. 2**SEQ ID NO: 2**

MVCEQPEVFASACALARAFPLFTHRSGASRRLEKKTVTVEFFLVGQDN
GPVEVSTLQCLANATDGVRLAARIVDTPCNEMNTDTFLEEINKVGKEL
GIIPTIIRDEELKTRGFGGIYGVGKAALHPPALAVLSHTPDGATQTIA
WVGKGIVYDTGGLSIKGKTTMPGMKRDCGAAAVLGAFRAAIKQGFKD
NLHAVFCLAENSVGP NATRPDDIHLLYSGKTVEINNTDAEGRVLADG
VSYACKDLGADIILDMATLTGAQGIATGKYHAAVLTNSAEWEAACVKA
GRKCGDLVHPLVYCP ELHFSEFTSAVADMKN SVADRDN SPSSCAGLFI
ASHIGFDWPGVWVHLDIAAPVHAGERATGFGVALLLALFGRASEDPLL
NLVSPLGCEVDVEEGDLGRDSKRRRLV

Fig. 3**SEQ ID NO: 3**

5' - TTTTCCTGGTGGGACAAGACA - 3'

Fig. 4**SEQ ID NO: 4**

5' - CATTCGCTAAGCACTGCAATG - 3'

Fig. 5

SEQ ID NO: 5

5' - CGGGCCGGTGGAGGTGTC - 3'